

SEQUENCE LISTING

<110> Kimble, Judith E
Bluelloch, Robert H

<120> Agent and Method for Modulating Cell Migration

5 <130> 960296.95386

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<150> 60/087170

<151> 1998-05-29

10 <150> 60/129023

<151> 1999-04-13

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<170> PatentIn Ver. 2.0

<210> 1

15 <211> 6659

<212> DNA

<213> Caenorhabditis elegans

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<221> CDS

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Ala Leu Ile Leu Leu Val Val Cys Leu Val Tyr Ala Leu Gln Ser Gly
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Ser Gly Thr Ile Ser Glu Phe Ser Ser Asp Val Leu Phe Ser Arg Ala
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5/11/1
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5 Gly Ile His Val Ile Asp Ser His His Ile Val Arg Arg Asp Ser Tyr
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10 Leu Gln Gly Val Ala Arg Asp Cys Gly His Ala Cys His Leu Arg Leu
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15 Arg Ser Asp Asp Ala Val Tyr Ile Val His Leu His Arg Trp Asn Gln
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5 Asp Trp Arg Ser Trp Gly Glu Cys Ser Arg Thr Cys Gly Gly Gly Val
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66650-66650

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SUB
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Ser Gln Thr Cys Glu Leu Asn Pro Cys Asp Ser Glu Phe Lys Trp Ser
1860 1865 1870

15 ttc gga cca tgg ggt gaa tgc tcg aaa aat tgc ggt caa ggt att cga 5664
Phe Gly Pro Trp Gly Glu Cys Ser Lys Asn Cys Gly Gln Gly Ile Arg
1875 1880 1885

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Arg Arg Arg Val Lys Cys Val Ala Asn Asp Gly Arg Arg Val Glu Arg
1890 1895 1900

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Val Lys Cys Thr Thr Lys Lys Pro Arg Arg Thr Gln Tyr Cys Phe Glu
1905 1910 1915 1920

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Arg Asn Cys Leu Pro Ser Thr Cys Gln Glu Leu Lys Ser Gln Asn Val
1925 1930 1935

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Lys Ala Lys Asp Gly Asn Tyr Thr Ile Leu Leu Asp Gly Phe Thr Ile
1940 1945 1950

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Glu Ile Tyr Cys His Arg Met Asn Ser Thr Ile Pro Lys Ala Tyr Leu
1955 1960 1965

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Asn Val Asn Pro Arg Thr Asn Phe Ala Glu Val Tyr Gly Lys Lys Leu
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 5 His Cys Ser Glu Asp Gly Asp Ala Ser Ala Gly Leu Thr Arg Phe Asn
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<212> PRT

<213> Caenorhabditis elegans

<400> 2

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Ser Gly Thr Ile Ser Glu Phe Ser Ser Asp Val Leu Phe Ser Arg Ala
35 40 45

15 Lys Tyr Ser Gly Val Pro Val His His Ser Arg Trp Arg Gln Asp Ala
50 55 60

Gly Ile His Val Ile Asp Ser His His Ile Val Arg Arg Asp Ser Tyr
65 70 75 80

20 Gly Arg Arg Gly Lys Arg Asp Val Thr Ser Thr Asp Arg Arg Arg Arg
85 90 95

Leu Gln Gly Val Ala Arg Asp Cys Gly His Ala Cys His Leu Arg Leu
100 105 110

Arg Ser Asp Asp Ala Val Tyr Ile Val His Leu His Arg Trp Asn Gln
115 120 125

25 Ile Pro Asp Ser His Asn Lys Ser Val Pro His Phe Ser Asn Ser Asn
130 135 140

Phe Ala Pro Met Val Leu Tyr Leu Asp Ser Glu Glu Glu Val Arg Gly
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30 Gly Met Ser Arg Thr Asp Pro Asp Cys Ile Tyr Arg Ala His Val Lys
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Gly Val His Gln His Ser Ile Val Asn Leu Cys Asp Ser Glu Asp Gly
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Leu Tyr Gly Met Leu Ala Leu Pro Ser Gly Ile His Thr Val Glu Pro
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5 Ile Ile Ser Gly Asn Gly Thr Glu His Asp Gly Ala Ser Arg His Arg
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Gln His Leu Val Arg Lys Phe Asp Pro Met His Phe Lys Ser Phe Asp
225 230 235 240

10 His Leu Asn Ser Thr Ser Val Asn Glu Thr Glu Thr Thr Val Ala Thr
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Trp Gln Asp Gln Trp Glu Asp Val Ile Glu Arg Lys Ala Arg Ser Arg
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Arg Ala Ala Asn Ser Trp Asp His Tyr Val Glu Val Leu Val Val Ala
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15 Asp Thr Lys Met Tyr Glu Tyr His Gly Arg Ser Leu Glu Asp Tyr Val
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Leu Thr Leu Phe Ser Thr Val Ala Ser Ile Tyr Arg His Gln Ser Leu
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20 Arg Ala Ser Ile Asn Val Val Val Val Lys Leu Ile Val Leu Lys Thr
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Glu Asn Ala Gly Pro Arg Ile Thr Gln Asn Ala Gln Gln Thr Leu Gln
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Asp Phe Cys Arg Trp Gln Gln Tyr Tyr Asn Asp Pro Asp Asp Ser Ser
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30 Met Cys Asp Met Gln Lys Ser Cys Ala Ile Ile Glu Asp Asn Gly Leu
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Ser Ala Ala Phe Thr Ile Ala His Glu Leu Gly His Val Phe Ser Ile
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 Pro His Asp Asp Glu Arg Lys Cys Ser Thr Tyr Met Pro Val Asn Lys
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 Trp Ser Trp Ser Pro Cys Ser Ala Gly Met Leu Glu Arg Phe Leu Glu
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 25 Gln Lys Gly Leu Arg Asp Cys Asp Ser Pro Lys Pro Arg Asn Gly Gly
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 Lys Tyr Cys Val Gly Gln Arg Glu Arg Tyr Arg Ser Cys Asn Thr Gln
 625 630 635 640
 30 Glu Cys Pro Trp Asp Thr Gln Pro Tyr Arg Glu Val Gln Cys Ser Glu
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Phe Asn Asn Lys Asp Ile Gly Ile Gln Gly Val Ala Ser Thr Asn Thr
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 His Trp Val Pro Lys Tyr Ala Asn Val Ala Pro Asn Glu Arg Cys Lys
 675 680 685
 5 Leu Tyr Cys Arg Leu Ser Gly Ser Ala Ala Phe Tyr Leu Leu Arg Asp
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 Lys Val Val Asp Gly Thr Pro Cys Asp Arg Asn Gly Asp Asp Ile Cys
 705 710 715 720
 Val Ala Gly Ala Cys Met Pro Ala Gly Cys Asp His Gln Leu His Ser
 10 725 730 735
 Thr Leu Arg Arg Asp Lys Cys Gly Val Cys Gly Gly Asp Asp Ser Ser
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Leu Asp Met Ser Thr His Arg Gln Ser His Asp Arg Asn Cys Gln Asn
915 920 925

5 Val Leu Lys Pro Lys Gln Ala Thr Arg Met Cys Asn Ile Asp Cys Ser
930 935 940

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10 Ser Gly Gln Lys Arg Gln Arg Val Ser Cys Val Lys Met Glu Gly Asp
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995 1000 1005

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 30 1845 1850 1855

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